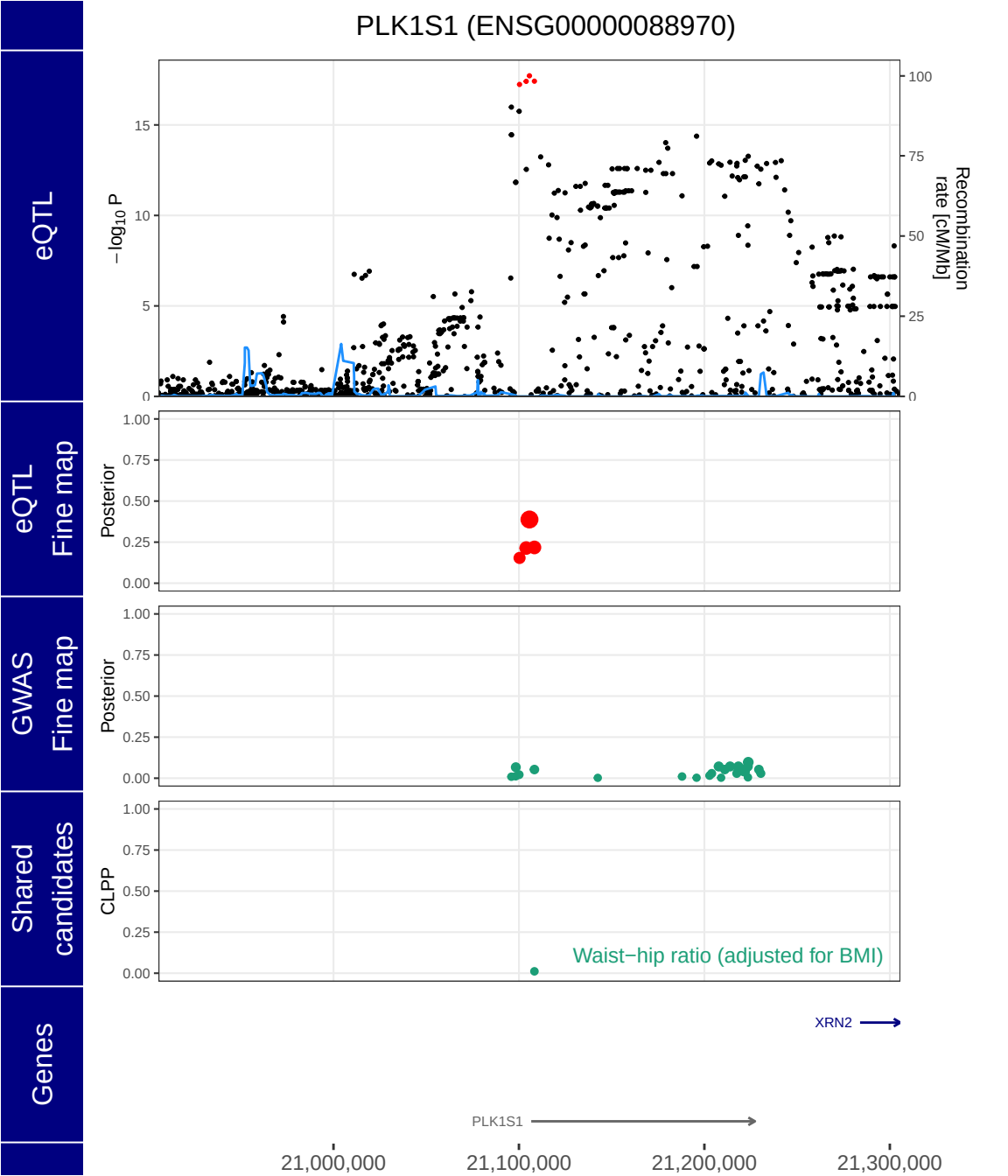
PLK1S1 (ENSG00000088970)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Waist−hip ratio (adjusted for BMI)
XRN2
PLK1S1
21,000,000
21,100,000
21,200,000
21,300,000